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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
09/580,110	05/30/2000	Thomas F. Mitts	00-142-US	1492

7590

12/10/2002

RAYMOND A. MILLER, ESQ.  
BENESCH, FRIEDLANDER, COPLAN & ARONOFF, LLP  
2300 BP TOWER  
200 PUBLIC SQUARE  
CLEVELAND, OH 44114-2378

EXAMINER

SHEINBERG, MONIKA B

ART UNIT

PAPER NUMBER

1634

DATE MAILED: 12/10/2002

14

Please find below and/or attached an Office communication concerning this application or proceeding.



UNITED STATES DEPARTMENT OF COMMERCE  
Patent and Trademark Office

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Washington, D.C. 20231

APPLICATION NO./ CONTROL NO.	FILING DATE	FIRST NAMED INVENTOR / PATENT IN REEXAMINATION	ATTORNEY DOCKET NO.
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EXAMINER
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ART UNIT	PAPER
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14

DATE MAILED:

**Please find below and/or attached an Office communication concerning this application or proceeding.**

**Commissioner of Patents and Trademarks**

**Sequence Compliance**

This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 CFR § 1.821(a)(1) and (a)(2). It is acknowledged that there is a Sequence Listing in computer readable format. Unfortunately, this application fails to comply with the requirements of 37 CFR § 1.821 through 1.825 because the reasons listed in the attached "Raw Sequence Listing Error Report". Thus, it is requested that the applicant send in another copy of the computer readable format. Please provide another statement as per 37 CFR § 1.821 (f) stating that the Paper Listing and the computer readable format are the same. Applicant is required to complete the response within a time limit of one month from the date of this letter or as extended as follows. **AN EXTENSION OF THIS TIME LIMIT MAY BE GRANTED UNDER EITHER 37 CFR § 1.136 (a) OR (b) UP TO A MAXIMUM OF SIX MONTHS.**

In addition, please note that claim 5 appears to be incomplete due to typographical errors. Applicant is requested to correct claim 5 to be in complete format.


**Inquiries**

Papers related to this application may be submitted to Technical Center 1600 by facsimile transmission. Papers should be faxed to Technical Center 1600 via the PTO Fax Center located in Crystal Mall 1. The faxing of such papers must conform with the notices published in the Official Gazette, 1096 OG 30 (November 15, 1988), 1156 OG 61 (November 16, 1993), and 1157 OG 94 (December 28, 1993) (See 37 CFR § 1.6(d)). The CM1 Fax Center number is (703) 308-4242.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Monika B. Sheinberg, whose telephone number is (703) 306-0511. The examiner can normally be reached on Monday-Friday from 1 P.M. to 8 P.M. If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Gary Jones, can be reached on (703) 308-1152.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to Patent Analyst, Chantae Dessau, whose telephone number is (703) 605-1237, or to the Technical Center receptionist whose telephone number is (703) 308-0196.

November 20, 2002  
Monika B. Sheinberg

  
W. Gary Jones  
Supervisory Patent Examiner  
Technology Center 1600

# NOTICE TO COMPLY WITH SEQUENCE RULES

Application No.

09/580110

Examiner

Monika Sheinberg

Applicant(s)

ts et al.

Art Unit

1634

## NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821-1.825 for the following reasons:

- ☒ 1. This application clearly fails to comply with the requirements of 37 CFR 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
- ☐ 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 CFR 1.821(c).
- ☐ 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 CFR 1.821(e).
- ☐ 4. A copy of the "Sequence Listing in computer readable form has been submitted. However the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached copy of the marked up "Raw Sequence Listing".
- ☐ 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable. A Substitute computer readable form must be submitted as required by 37 CFR 1.825(d).
- ☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e).
- ☒ 7. Other: Please see the attached "Raw Sequence Listing Error Report"

### Applicant must provide:

- ☐ An initial or ☐ A substitute computer readable form copy of the Sequence Listing.
- ☐ An initial or ☐ A Substitute paper copy of the Sequence Listing as well as an amendment directing its entry into the specification.
- ☒ A statement that the content of the paper and computer readable copies are the same, and, where applicable, include no new matter, as required by 37 CFR 1.821(e), (f), or (g) or 1.825(b) or (d).

### FOR QUESTIONS PLEASE CONTACT:

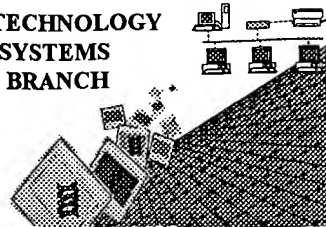
Rules Interpretation (703) 308-4216  
CRF Submission Help (703) 308 4212  
PatentIn software help (703) 308 6856

**PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE**

*Tamela Taylor*

**RAW SEQUENCE LISTING**  
**ERROR REPORT**

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



1694

**RECEIVED**

OCT 30 2002

TECH CENTER 1600/2900

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/580,110P

Source: 1600 Rush

Date Processed by STIC: 10/24/2002

#13  
CO  
11/13/02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER**  
**VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND  
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efb/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised Q1/29/2002

RECEIVED

OCT 30 2002

TECH CENTER 1600/2900

## Raw Sequence Listing Error Summary

ERROR DETECTEDSUGGESTED CORRECTIONSERIAL NUMBER: 09/580,1100

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1      Wrapped Nucleics  
    Wrapped Aminos  
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2      Invalid Line Length  
The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3      Misaligned Amino  
    Numbering  
The numbering under each 5<sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4      Non-ASCII  
The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5      Variable Length  
Sequence(s)      contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6      PatentIn 2.0  
    "bug"  
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)     . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7      Skipped Sequences  
    (OLD RULES)  
Sequence(s)      missing. If intentional, please insert the following lines for each skipped sequence:  
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
This sequence is intentionally skipped  
  
Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8      Skipped Sequences  
    (NEW RULES)  
Sequence(s)      missing. If intentional, please insert the following lines for each skipped sequence.  
<210> sequence id number  
<400> sequence id number  
000
- 9      Use of n's or Xaa's  
    (NEW RULES)  
Use of n's and/or Xaa's have been detected in the Sequence Listing.  
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  
In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10    Invalid <213>  
    Response  
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11      Use of <220>  
Sequence(s)      missing the <220> "Feature" and associated numeric identifiers and responses.  
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12      PatentIn 2.0  
    "bug"  
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13      Misuse of n  
n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

J.  
Taylor

1600

## RAW SEQUENCE LISTING

DATE: 10/24/2002

PATENT APPLICATION: US/09/580,110D

TIME: 13:19:15

Input Set : A:\09-580110.txt

Output Set: N:\CRF4\10242002\I580110D.raw

3 <110> APPLICANT: MITTS, THOMAS F.  
 4 SANDBERG, LAWRENCE B.  
 6 <120> TITLE OF INVENTION: ELASTIN PEPTIDE ANALOGS AND USES OF SAME IN COMBINATION  
 7 WITH SKIN ENHANCING AGENTS  
 9 <130> FILE REFERENCE: 00-142-US  
 11 <140> CURRENT APPLICATION NUMBER: 09/580,110D  
 12 <141> CURRENT FILING DATE: 2000-05-30  
 14 <160> NUMBER OF SEQ ID NOS: 75  
 16 <170> SOFTWARE: PatentIn Ver. 2.1  
 18 <210> SEQ ID NO: 1  
 19 <211> LENGTH: 3  
 20 <212> TYPE: PRT  
 21 <213> ORGANISM: mammalian  
 23 <400> SEQUENCE: 1  
 24 Ala Val Gly  
 25 1  
 28 <210> SEQ ID NO: 2  
 29 <211> LENGTH: 4  
 30 <212> TYPE: PRT  
 31 <213> ORGANISM: mammalian  
 33 <400> SEQUENCE: 2  
 34 Val Gly Ala Gly  
 35 1  
 38 <210> SEQ ID NO: 3  
 39 <211> LENGTH: 3  
 40 <212> TYPE: PRT  
 41 <213> ORGANISM: mammalian  
 43 <400> SEQUENCE: 3  
 44 Ile Gly Gly  
 45 1  
 48 <210> SEQ ID NO: 4  
 49 <211> LENGTH: 2  
 50 <212> TYPE: PRT  
 51 <213> ORGANISM: mammalian  
 53 <400> SEQUENCE: 4  
 54 Leu Gly  
 55 1  
 58 <210> SEQ ID NO: 5  
 59 <211> LENGTH: 4  
 60 <212> TYPE: PRT  
 61 <213> ORGANISM: mammalian  
 63 <400> SEQUENCE: 5  
 64 Ile Gly Ala Gly

*invalid (global error)*  
*see item 10 on*  
*Erra summary*  
*sheet*

Does Not Comply  
 Corrected Diskette Needed

*pg 1-6*

## RAW SEQUENCE LISTING

DATE: 10/24/2002

PATENT APPLICATION: US/09/580,110D

TIME: 13:19:15

Input Set : A:\09-580110.txt

Output Set: N:\CRF4\10242002\I580110D.raw

65 1  
68 <210> SEQ ID NO: 6  
69 <211> LENGTH: 3  
70 <212> TYPE: PRT  
71 <213> ORGANISM: mammalian  
73 <400> SEQUENCE: 6  
74 Leu Gly Gly  
75 1  
78 <210> SEQ ID NO: 7  
79 <211> LENGTH: 4  
80 <212> TYPE: PRT  
81 <213> ORGANISM: mammalian  
83 <400> SEQUENCE: 7  
84 Val Ala Pro Gly  
85 1  
88 <210> SEQ ID NO: 8  
89 <211> LENGTH: 4  
90 <212> TYPE: PRT  
91 <213> ORGANISM: mammalian  
93 <400> SEQUENCE: 8  
94 Leu Gly Pro Gly  
95 1  
98 <210> SEQ ID NO: 9  
99 <211> LENGTH: 4  
100 <212> TYPE: PRT  
101 <213> ORGANISM: mammalian  
103 <400> SEQUENCE: 9  
104 Leu Gly Ala Gly  
105 1  
108 <210> SEQ ID NO: 10  
109 <211> LENGTH: 4  
110 <212> TYPE: PRT  
111 <213> ORGANISM: mammalian  
113 <400> SEQUENCE: 10  
114 Val Gly Pro Gly  
115 1  
118 <210> SEQ ID NO: 11  
119 <211> LENGTH: 4  
120 <212> TYPE: PRT  
121 <213> ORGANISM: mammalian  
123 <400> SEQUENCE: 11  
124 Phe Gly Pro Gly  
125 1  
128 <210> SEQ ID NO: 12  
129 <211> LENGTH: 4  
130 <212> TYPE: PRT  
131 <213> ORGANISM: mammalian  
133 <400> SEQUENCE: 12  
134 Val Gly Pro Gln

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/580,110D

DATE: 10/24/2002

TIME: 13:19:15

Input Set : A:\09-580110.txt

Output Set: N:\CRF4\10242002\I580110D.raw

135 1  
138 <210> SEQ ID NO: 13  
139 <211> LENGTH: 3  
140 <212> TYPE: PRT  
141 <213> ORGANISM: mammalian  
143 <400> SEQUENCE: 13  
144 Leu Gly Ala  
145 1  
148 <210> SEQ ID NO: 14  
149 <211> LENGTH: 4  
150 <212> TYPE: PRT  
151 <213> ORGANISM: mammalian  
153 <400> SEQUENCE: 14  
154 Val Gly Pro Ala  
155 1  
158 <210> SEQ ID NO: 15  
159 <211> LENGTH: 4  
160 <212> TYPE: PRT  
161 <213> ORGANISM: mammalian  
163 <400> SEQUENCE: 15  
164 Val Val Pro Gly  
165 1  
168 <210> SEQ ID NO: 16  
169 <211> LENGTH: 4  
170 <212> TYPE: PRT  
171 <213> ORGANISM: mammalian  
173 <400> SEQUENCE: 16  
174 Ala Val Pro Gly  
175 1  
178 <210> SEQ ID NO: 17  
179 <211> LENGTH: 4  
180 <212> TYPE: PRT  
181 <213> ORGANISM: mammalian  
183 <400> SEQUENCE: 17  
184 Val Val Pro Gln  
185 1  
188 <210> SEQ ID NO: 18  
189 <211> LENGTH: 6  
190 <212> TYPE: PRT  
191 <213> ORGANISM: mammalian  
193 <400> SEQUENCE: 18  
194 Val Ala Ala Arg Pro Gly  
195 1 5  
198 <210> SEQ ID NO: 19  
199 <211> LENGTH: 7  
200 <212> TYPE: PRT  
201 <213> ORGANISM: mammalian  
203 <400> SEQUENCE: 19  
204 Leu Gly Ala Gly Gly Ala Gly



## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/580,110D

DATE: 10/24/2002

TIME: 13:19:15

Input Set : A:\09-580110.txt

Output Set: N:\CRF4\10242002\I580110D.raw

205 1 5  
208 <210> SEQ ID NO: 20  
209 <211> LENGTH: 4  
210 <212> TYPE: PRT  
211 <213> ORGANISM: mammalian  
213 <400> SEQUENCE: 20  
214 Ala Ile Pro Gly  
215 1  
218 <210> SEQ ID NO: 21  
219 <211> LENGTH: 5  
220 <212> TYPE: PRT  
221 <213> ORGANISM: mammalian  
223 <400> SEQUENCE: 21  
224 Leu Gly Pro Gly Gly  
225 1 5  
228 <210> SEQ ID NO: 22  
229 <211> LENGTH: 5  
230 <212> TYPE: PRT  
231 <213> ORGANISM: mammalian  
233 <400> SEQUENCE: 22  
234 Ala Ala Ala Gln Ala  
235 1 5  
238 <210> SEQ ID NO: 23  
239 <211> LENGTH: 5  
240 <212> TYPE: PRT  
241 <213> ORGANISM: mammalian  
243 <220> FEATURE:  
244 <221> NAME/KEY: MOD\_RES  
245 <222> LOCATION: (4)  
247 <400> SEQUENCE: 23  
W--> 248 Val Gly Val Xaa Gly  
249 1 5  
252 <210> SEQ ID NO: 24  
253 <211> LENGTH: 5  
254 <212> TYPE: PRT  
255 <213> ORGANISM: mammalian  
257 <400> SEQUENCE: 24  
258 Val Tyr Pro Gly Gly  
259 1 5  
262 <210> SEQ ID NO: 25  
263 <211> LENGTH: 6  
264 <212> TYPE: PRT  
265 <213> ORGANISM: mammalian  
267 <400> SEQUENCE: 25  
268 Ile Gly Gly Val Gly Gly  
269 1 5  
272 <210> SEQ ID NO: 26  
273 <211> LENGTH: 6  
274 <212> TYPE: PRT

> <223> give explanation of Xaa (see p. 6)  
for error  
explanation

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/580,110D

DATE: 10/24/2002

TIME: 13:19:15

Input Set : A:\09-580110.txt

Output Set: N:\CRF4\10242002\I580110D.raw

275 &lt;213&gt; ORGANISM: mammalian

277 &lt;400&gt; SEQUENCE: 26

278 Val Ala Pro Gly Val Gly

279 1 5

282 &lt;210&gt; SEQ ID NO: 27

283 &lt;211&gt; LENGTH: 5

284 &lt;212&gt; TYPE: PRT

285 &lt;213&gt; ORGANISM: mammalian

287 &lt;400&gt; SEQUENCE: 27

288 Leu Gly Val Gly Gly

289 1 5

292 &lt;210&gt; SEQ ID NO: 28

293 &lt;211&gt; LENGTH: 4

294 &lt;212&gt; TYPE: PRT

295 &lt;213&gt; ORGANISM: mammalian

297 &lt;400&gt; SEQUENCE: 28

298 Leu Val Pro Gly

299 1

302 &lt;210&gt; SEQ ID NO: 29

303 &lt;211&gt; LENGTH: 5

304 &lt;212&gt; TYPE: PRT

305 &lt;213&gt; ORGANISM: mammalian

307 &lt;400&gt; SEQUENCE: 29

308 Phe Arg Ala Ala Ala

309 1 5

312 &lt;210&gt; SEQ ID NO: 30

313 &lt;211&gt; LENGTH: 6

314 &lt;212&gt; TYPE: PRT

315 &lt;213&gt; ORGANISM: mammalian

317 &lt;400&gt; SEQUENCE: 30

318 Val Gly Gly Val Pro Gly

319 1 5

322 &lt;210&gt; SEQ ID NO: 31

323 &lt;211&gt; LENGTH: 5

324 &lt;212&gt; TYPE: PRT

325 &lt;213&gt; ORGANISM: mammalian

327 &lt;400&gt; SEQUENCE: 31

328 Phe Gly Pro Gly Gly

329 1 5

332 &lt;210&gt; SEQ ID NO: 32

333 &lt;211&gt; LENGTH: 5

334 &lt;212&gt; TYPE: PRT

335 &lt;213&gt; ORGANISM: mammalian

337 &lt;400&gt; SEQUENCE: 32

338 Val Gly Val Pro Gly

339 1 5

342 &lt;210&gt; SEQ ID NO: 33

343 &lt;211&gt; LENGTH: 6

344 &lt;212&gt; TYPE: PRT

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/580,110D

DATE: 10/24/2002  
TIME: 13:19:16

Input Set : A:\09-580110.txt  
Output Set: N:\CRF4\10242002\I580110D.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:23; Xaa Pos. 4  
Seq#:34; Xaa Pos. 4

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/580,110D

DATE: 10/24/2002

TIME: 13:19:16

Input Set : A:\09-580110.txt

Output Set: N:\CRF4\10242002\I580110D.raw

L:248 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:23  
L:248 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0  
L:362 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:34  
L:362 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0